Marked-up Version of Proposed Amendments

Claim 6:

A mutant of any one of claims 1-7 claim 4 or claim 5, which comprises the sequence with PRPRSFSAPFSSQDS (SEQ ID No. 37), or a sequence substantially homologous thereto.

Claim 7:

A mutant of any one of claims 1-7 any one of claims 4 to 6 which comprises the sequence with MLRALNRLAARPGGQPPTLLLLPVRGPRPRSFSAPFSSQDS (SEQ ID No. 30), or a sequence substantially homologous thereto.

Claim 10:

A nucleic acid of any one of claims 1-7 claim 9, comprising the sequence with ATGTTGCGGGCTTTGAACCGCCTGGCCGCGCGGCCCGGGGGCCAGCCCCCAACCCT GCTCCTTCTGCCCGTGCGCGGCCCACGGCCCCGCTCATTCTCGGCTCCTTTTTCCTCG CAGGATAGC (SEQ ID No. 31), or an equivalent sequence which encodes the same polypeptide having regard to the degeneracy of the nucleic acid code, or a sequence substantially homologous thereto.

Page 6, fourth full paragraph

Figure 5 shows a point mutation in the cVO14 construct (SEQ ID NO: 36) in a conserved region 5' to the OT gene. A conserved G residue is substituted with an A residue in the construct compared to the sequence from for Mouse (SEQ ID NO: 32), Human (SEQ ID NO: 33), Cow (SEQ ID NO: 34), and Rat (SEQ ID NO: 35).